

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on Apr 16, 2025

Kaptive

RRID:SCR_024046

Type: Tool

Proper Citation

Kaptive (RRID:SCR_024046)

Resource Information

URL: <https://github.com/klebgenomics/Kaptive>

Proper Citation: Kaptive (RRID:SCR_024046)

Description: Software tool to report information about surface polysaccharide loci for *Klebsiella pneumoniae* species complex and *Acinetobacter baumannii* genome assemblies.

Synonyms: kaptive

Resource Type: software resource, software application

Defining Citation: [PMID:28348840](https://pubmed.ncbi.nlm.nih.gov/28348840/)

Keywords: report information, surface polysaccharide loci, genome assemblies,

Funding:

Availability: Free, Available for download, Freely available,

Resource Name: Kaptive

Resource ID: SCR_024046

Alternate IDs: OMICS_27932

Alternate URLs: <https://sources.debian.org/src/kaptive/>

License: GPL-3.0 license

Record Creation Time: 20230824T050211+0000

Record Last Update: 20250416T063952+0000

Ratings and Alerts

No rating or validation information has been found for Kaptive.

No alerts have been found for Kaptive.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Boeckaerts D, et al. (2024) Prediction of Klebsiella phage-host specificity at the strain level. *Nature communications*, 15(1), 4355.

Shpirt AM, et al. (2024) An *Acinetobacter baumannii* nasal carriage isolate recovered from an asymptomatic patient in Vietnam is extensively antibiotic resistant and produces a rare K71 type capsule. *Microbiology spectrum*, 12(12), e0183824.

Feng Y, et al. (2024) Population genomics uncovers global distribution, antimicrobial resistance, and virulence genes of the opportunistic pathogen *Klebsiella aerogenes*. *Cell reports*, 43(8), 114602.

Slarve MJ, et al. (2024) Clinical assays rapidly predict bacterial susceptibility to monoclonal antibody therapy. *JCI insight*, 9(2).

Amadesi S, et al. (2024) Complete Genome Sequence of a *Klebsiella pneumoniae* Strain Carrying Novel Variant blaKPC-203, Cross-Resistant to Ceftazidime/Avibactam and Cefiderocol, but Susceptible to Carbapenems, Isolated in Italy, 2023. *Pathogens (Basel, Switzerland)*, 13(6).

Yang Y, et al. (2024) Large-scale genomic survey with deep learning-based method reveals strain-level phage specificity determinants. *GigaScience*, 13.